

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasm mem
P25425	PO2F1_MOUSE	Mus musculus	POU domain, class 2, transcription factor 1	51.997308	T256	T271;T277;S284;S388;S451	23580612;36084651;36852467;40128797;40997131	MNNPSETNKSSMESEDASTGTQTN GLDFQKQPVVGGAISTAQAQAF LG HLHQVQLAGTSLQAAAQSLNVQSKS SEESGDSQQSSQPSSQPPSVQSAIPQ TQLMLAGGQITGLTLTPAQQLLLQ QAQAQAQLLAAAQQHSASQQHSA AGATISASAATPMTQIPLSQPIQIAD LQQLQQLQQQNLNLQQFVLVHPTT NLQPAQFIISQTPQGGQGLLQAQNL LTQLPQQSQANLLQPQPSITLTSQPT TPTRTIAAASVQTLPQSQSTPKRIDTP SLEEPSDLEELEQFAKTFKQRRIKLG FTQGDVGLAMGKLYGNDFSQTTISR FEALNLSFKNMCKLKPILLEKWLND AENLSSDSTASSPSALNSPGLGAEG LNRRRKKRTSIETNIRVALEKSFME NQKPTSEDIILIAEQLNMEKEVIRV WFCNRRQKEKRINPPSSGGTSSSPI KAIFPSPASLVATTPLVTSSTATTTLT VNPVPLTSAAVTNLSLTDQLRRG CSWEVLRSLPDRVTTTAGTTDSTSN NNTATVISTAPPASSAVTSPSLSPSPS ASASTSEASSASETNTTQTTSTPLPS PLGASQVMVTTPLQLTAAAAALQGAA QLPANASLAAMAAAAGLSPGLMAPS QFAAGGALLSLSPGTLGSALSPALM SNSTLATIQALASSGSLPITSLDATGN LVFANAGGAPNIVTAPLFLNPQNL SL LTSNPVSLVSAAAASTGNSAPTASLH ASSTSTESIQSSLFTVASASGPASTTT AASKAQ	True	False	1.238	5.0	1.402	2.653	0.893	1.436